



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/966955

Source: OIE

Date Processed by STIC: 10/18/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/966,955

DATE: 10/18/2001

TIME: 09:04:19

Input Set : A:\3053us11.app

Output Set: N:\CRF3\10182001\I966955.raw

3 <110> APPLICANT: Perez-Villar, Juan J.

4 Chang, Han

5 Yang, Wen-Pin

6 Wu, Yuli

7 Whitney, Gena S.

8 Kanner, Steven B.

**Does Not Comply
Corrected Diskette Needed**

10 <120> TITLE OF INVENTION: Identification and Cloning of a Full-length Human

11 Clnk-related Gene, MIST (Mast Cell Immunoreceptor

12 Signal Transducer)

14 <130> FILE REFERENCE: 3053-4113US1

C--> 16 <140> CURRENT APPLICATION NUMBER: US/09/966,955

17 <141> CURRENT FILING DATE: 2001-09-28

19 <150> PRIOR APPLICATION NUMBER: 60/237030

20 <151> PRIOR FILING DATE: 2000-09-29

22 <160> NUMBER OF SEQ ID NOS: 52

24 <170> SOFTWARE: PatentIn Ver. 2.1

*use correct current application
number*

ERRORED SEQUENCES

462 <210> SEQ ID NO: 8

463 <211> LENGTH: 21

464 <212> TYPE: DNA

465 <213> ORGANISM: HUMAN

467 <220> FEATURE:

468 <223> OTHER INFORMATION: PRIMER PY474

470 <400> SEQUENCE: 8

E--> 471 ~~tggtacattg gagaatacag~~Errored:*Input: 21 nucleotides on field 211 Sequence 8
Found: 20 nucleotides on actual sequence***Does Not Comply
Corrected Diskette Needed**

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/966,955

DATE: 10/18/2001

TIME: 09:04:20

Input Set : A:\3053us11.app

Output Set: N:\CRF3\10182001\I966955.raw

L:16 M:270 C: Current Application Number differs, Wrong Format

L:471 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:20 SEQ:8